

BRS L1	2 20030082800.pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L2	2 6054299.pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L3	2 5436141.pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L4	120 ssDNA with expression	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L5	3734 ((single adj strand adj dna) or ssDNA with expression	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L6	159 ((single adj strand adj dna) or ssDNA) with expression	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L7	47 reverse adj transcriptase and l6	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L8	4 reverse adj transcriptase and l6 and retron	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT

L6 ANSWER 1 OF 1 REGISTRY COPYRIGHT 2007 ACS on STN
RN 336761-66-1 REGISTRY
CN DNA (*Corynebacterium glutamicum* strain ATCC13032 clone
EP1108790-Sequence-3471 gene) (9CI) (CA INDEX NAME)
OTHER NAMES:
CN 3470: PN: EP1108790 SEQID: 3471 claimed DNA
FS NUCLEIC ACID SEQUENCE
SQL 1230
NA 282 a 346 c 352 q 250 t

PATENT ANNOTATIONS (PNTE):

Sequence | Patent
Source | Reference

Not Given|EP1108790
|claimed SEQID
|3471

SEQ	1 atgcttaggca tgcttcgatg gactacagca ggtgaatccc acggccaggc 51 gcttatacgcc acgggtgaac acatgccagc aggctgccc gtgactaaag 101 atgaggtctc gtatcaattt ggcgcggcgc gcctggata tggctcgccgc 151 gctcgcatga agtttgagca agacgcgtt accttcctca cccgcatccg 201 ccacggccctc actttggta gccccatctc aatcatgatc ggcaacactg 251 agtgggataaa gtggaccacc atcatgtcct ctgacgctt ggacatggaa 301 gacccagata acgttgcggc gatgtcttcg gttcggggcg caaaactgac 351 tcgtccgcgt ccagggccacg ctgattacgc aggcatgetc aagtacggat 401 tcgatgtatgc ccgcaacgtt ctggagcgtt cttcagcccg tgagacggca 451 gcacgcgtgg cagcagcaac cgttgcgcgt tccttcctgc gtgaaacctt 501 gggcgtggaa gtgcttccc acgttaatttc cattggtgcg ===== ===== 551 acactggcgc ggagccaacc tttgcagata ttcaagcaat cgatgattcc 601 ccagttcggt cattcggtaa agacgctgaa gaatccatga tcgcggaaat 651 cgaggccgca aagaagccg gcgataccct cgggtggcatc gtggaaagtga 701 ttgttgaagg cctgcccattc gtgttggct cacacatttc tggcgaagat 751 cgcctcgatg cgcagatcgc agctgcactc atggcattc aggcacatcaa 801 gggcgtggaa atcgggtgacg gtttgcagaaga agctcgtcga cgtggctccg 851 aagcccacga tgaagtgttc ctggatgaca acggcgtata cgcacacacc 901 aaccgtgcag gtggcctcga aaggcggcatg accaacggtg aaaccctgcg 951 cgttcgtgtc ggcataaggc caatttctac tgtgcctcgc gcccgtaaaa 1001 ccattgatata gaaaaacggc aaggcagcaa ccggaatcca ccagcgttcc 1051 gacgtgtgcg ctgttccagc cgccgggtgc gttgcagaag caatggtcac 1101 cctggttctc gccccgcgcag tcctgcagaa attcggcgtt gactccotga 1151 gcgaaacccaa gagcaacatt gacacctacc tcaaaaacat tgaggaacgaa 1201 atgaaattcg aagggtttaga ggtatggagcg
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HITS AT: 534-546

****RELATED SEQUENCES AVAILABLE WITH SEQLINK****

MF Unspecified

CI MAN

SR CA

LC STN Files: CA, CAPLUS, TOXCENTER, USPATFULL

DT.CA CAplus document type: Patent

RL.P Roles from patents: BIOL (Biological study); OCCU (Occurrence); PRP (Properties)

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE, SCISEARCH, WPIDS' ENTERED AT 14:15:43 ON
16 APR 2007

L1 9094 S (SINGLE(W) STRAND) (2W) DNA OR SSDNA WITH EXPRESS?
L2 11010 S (SINGLE(W) STRAND) (2W) DNA OR SSDNA(L) EXPRESS?
L3 2957 S ((SINGLE(W) STRAND) (2W) DNA OR SSDNA) (L) EXPRESS?
L4 0 S L3 AND REVERSE(W) TRANSSCRIPTASE
L5 85 S L3 AND REVERSE(W) TRANSCRIPTASE
L6 67 DUPLICATE REMOVE L5 (18 DUPLICATES REMOVED)

(FILE 'HOME' ENTERED AT 16:47:51 ON 16 AUG 2007)

FILE 'REGISTRY' ENTERED AT 16:48:10 ON 16 AUG 2007

L1 3 S TGGTGCCTCCGAG/SQEN
L2 3 SEQLINK EXACT L1

FILE 'CAPPLUS' ENTERED AT 16:50:44 ON 16 AUG 2007

L3 3 S L2
L4 3 S L1
L5 0 S L4 NOT L3

FILE 'REGISTRY' ENTERED AT 16:59:11 ON 16 AUG 2007

L6 3 S GTTTCGAAGGCTAGCTACAAACGATCATCCAG/SQEN

FILE 'CAPPLUS' ENTERED AT 16:59:59 ON 16 AUG 2007

L7 3 S L6
L8 3 S L7 NOT LL3
L9 1 S L7 NOT L3

FILE 'REGISTRY' ENTERED AT 17:03:55 ON 16 AUG 2007

L10 29 S GGCTAGCTACAAACGA/SQEN

SEQID 7

FILE 'REGISTRY' ENTERED AT 17:04:24 ON 16 AUG 2007

L11 29 S L10 NOT L3 NOT L7